Table 2. Pairs of primer sequences used for SSCP analysis of *BLM*.

Name	Forward sequence[a]	SEQ. ID. NO.	Reverse sequence[a]	SEQ. ID. NO.	Product length (bp)
C1-B	GGATCCTGGTTCCGTCCGC	(10)	GAGGTTCACTGAAGGAAAAGTC	(26)	269
C1-A	CAACTAGAACGTCACTCAGCC	(11)	GAAGTCCTTGACCCTTGCTG	(27)	233
C1-1	GACTTTTCCTTCAGTGAACCTC	(12)	GGGATTTCTTTACAGTTGGTGTG	(28)	186
C1-2	CCAGATTTCTTGCAGACTCCG	(13)	CTCTTACAAAGTGACTTTGGGG	(29)	213
C1-3	CTTTAAGTACCATCAATGATTGGG	(14)	CCTCAGTCAAATCTATTTGCTCG	(30)	227
C1-4	GAGTAAGCACTGCTCAGAAATC	(15)	GCTTAACCATTCTGAGTCATCC	(31)	160
C1-5	CGAGCAAATAGATTTGACTGAGG	(16)	CAATACATGGAACTTCTCAGTTG	(32)	223
C1-6	GAAGATCTCAGGAAAGTGAC	(17)	CGTACTAAGGCATTTTGAAGAGG	(33)	215
C1-7	CAACTGAGAAAGTTCCATGTATTG	(18)	CACAGTCTGTGCTGGTTTCTG	(34)	239
C1-9	CTATTCCTGATGATAAACTGAAAC	(19)	CCTTCATAGAATTCCCTGTAGG	(35)	200
C1-10	GTGGAGAGATACAGGCCTGATTC	(20)	GTGTTCAGCCCAGTTGCTAC	(36)	244
C1-11	CAGGATTTCTCTGCCACCAGG	(21)	GCAGTATGTTTATTCTGATCTTTC	(37)	183
C1-12	CAGGAAAATGTTCTCACAAGCAC	(22)	CCTTGATGGGGTTGATAGGCAG	(38)	203
C1-13	CAGCCAGCAAATCTTCCACAG	(23)	CGCTCATCATGTTTCAGATTTCTGG	(39)	204
C1-14	GAATTATACTGACAAGTCAGCAC	(24)	GATCTACGATAAGTGATCTCAAG	(40)	295
C1-15	CTCCTGGGGTTCACTGTTGTC	(25)	GAGTCTGTTACTTGCACAGATC	(41)	211
C1-16	CAATCATAAACTTCTATATGTCAC	(42)	GCCATCACCGGAACACAGAGG	(57)	207
C1-17	GTGGGGACATGATTTTCGTCAAG	(43)	GATTATGTCTCGTTAAAGCTCATG	(58)	175
C1-18	GACATCCTGACTCAGCTGAAG	(44)	CGTGTCAGCCCATGGTGTCAC	(59)	203
C1-19	GCACCACCCATATGATTCAGG	(45)	CAGATAACC TGA CAGCCATCC	(60)	179
C1-20	GATGAAGTGCAGCAGAAGTGG	(46)	CAGTCTGGTCACATCATGATAG	(61)	221
C1-21	GCAGAGACTGGAAGAGATGGG	(47)	GCTCGTATTCTCCCTGCATTCCG	(62)	188
C1-22	GTATAGCATGGTACATTACTGTG	(48)	CCTTGTGATGAACTATGTTCTTG	(63)	228
C1-23	GACTGACGATGTGAAAAGTATTG	(49)	CCAAAATCTGTGAACTATCAGC	(64)	235
C1-24	CCAGTCAGGTATATTTGGAAAAG	(50)	GGAATTTTCTTGTTTCCATAAAGTC	(65)	206
C1-25	CGATCGCTTATGTGATGCTCG	(51)	CAAGCTTCTTGAGAGTGACGG	(66)	248
C1-26	GAACTTACAGAAGTCTGCAAATC	(52)	GATGTCCATTCAGAGTATTTCTG	(67)	208
C1-27	GGTGTTACTGAAGAACAAACTGG	(53)	GGGTATTTCCTCGTCAAGCTC	(68)	168
C1-28	GGATAAGCCTGTCCAGCAGC	(54)	CCTAGATATCTTTCTACATGTGG	(69)	214
C1-29	GCTTCCAGTGGTTCCAAGGC	(55)	GTTATGAGAATGCATATGAAGGC	(70)	204
C1-30	CTCAAGCGACATCAGGAGCC	(56)	CAAGAATAACAGCTTTATAGTCAC	(71)	178

[a] 5' to 3'